

0916423.093098

HU:MGSDVRDLNALLPAVPSLGGGGGCALPVSGAAQWAPVLDFAAPPASAYGSL
MO:MGSDVRDLNALLPAVSSLGGGGGCGLPVSGAAQWAPVLDFAAPPASAYGSL

HU:GGPAPPPAPPPPPPPPPHSHFIKQEPSWGGAEPEEQCLSAFTVHFSGQFTGTAG
MO:GGPAPPPAPPPPPPPPPHSHFIKQEPSWGGAEPEEQCLSAFTLHFSGQFTGTAG

HU:ACRYGPFPPPPPSQASSGQARMFPNAPYLPSCLESQPAIRNQGYSTVTFDGTFS
MO:ACRYGPFPPPPPSQASSGQARMFPNAPYLPSCLESQPTIRNQGYSTVTFDGAPS

HU:YGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVYGCHTPTDSCGT
MO:YGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVYGCHTPTDSCGT

HU:SQALLLRTPYSSDNLQMTSQLECMTNQMNLGATLKGVAAGSSSSSVKWTE
MO:SQALLLRTPYSSDNLQMTSQLECMTNQMNLGATLKGMAAGSSSSSVKWTE

HU:GQSNHSTGYESDNHTTPILCGAQYRIHTGVFRGIQDVRRVPGVAPTLVRSAS
MO:GQSNHGIGYESDNHTAPILCGAQYRIHTGVFRGIQDVRRVSGVAPTLVRSAS

HU:ETSEKRPFMCAYPGCNKRYFKLSHLQMHRSKHTGEKPYQCDFKDCERRFSR
MO:ETSEKRPFMCAYPGCNKRYFKLSHLQMHRSKHTGEKPYQCDFKDCERRFSR

HU:SDQLKRHQRRHTGVKPFQCKTCQKFSRSDHLKTHTRTHTGKTSEKPFSCR
MO:SDQLKRHQRRHTGVKPFQCKTCQKFSRSDHLKTHTRTHTGKTSEKPFSCR

HU:WPSCQKKFARSDELVRHHNMHQRMNTKLQAL
MO:WHSCQKKFARSDELVRHHNMHQRMNTKLHVAL

FIG. 1

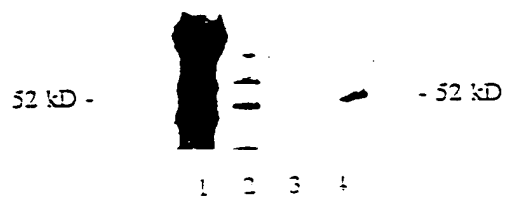


FIG. 2

0916423 093060
093060 093060

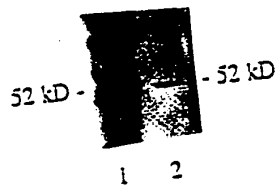


FIG. 3

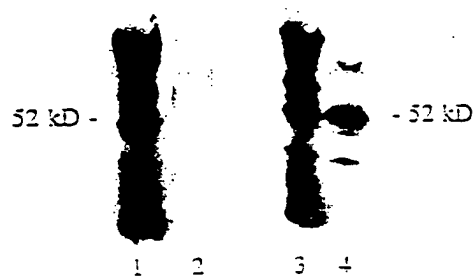


FIG. 4

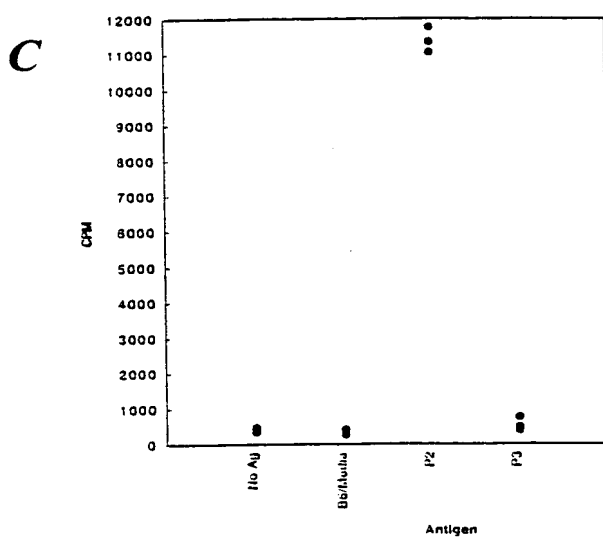
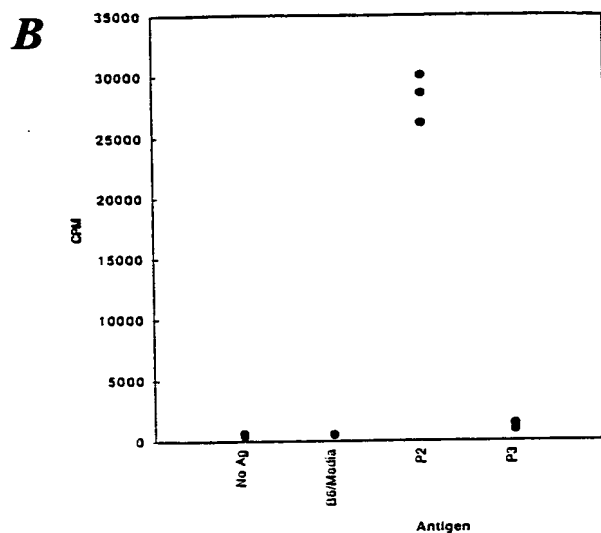
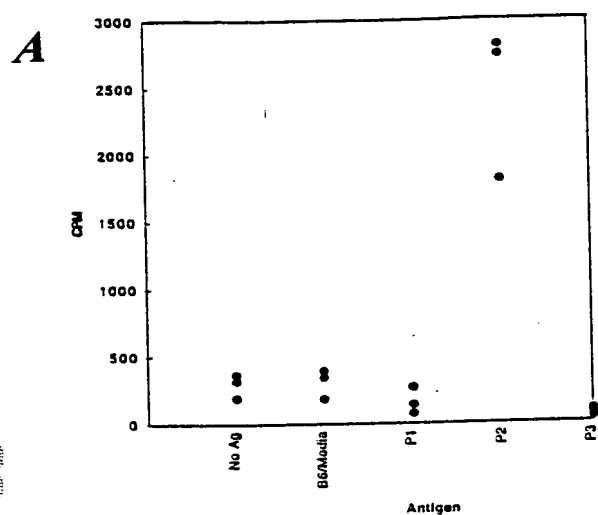
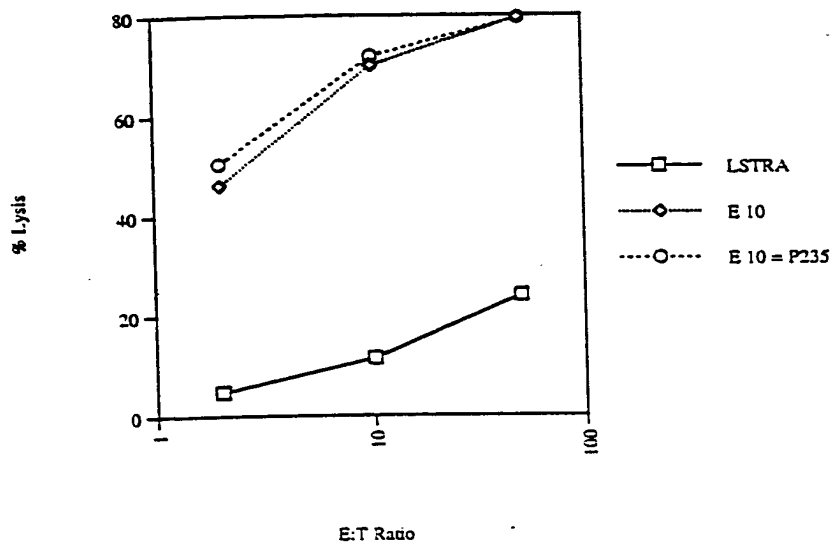


FIG. 5A-5C

A



B

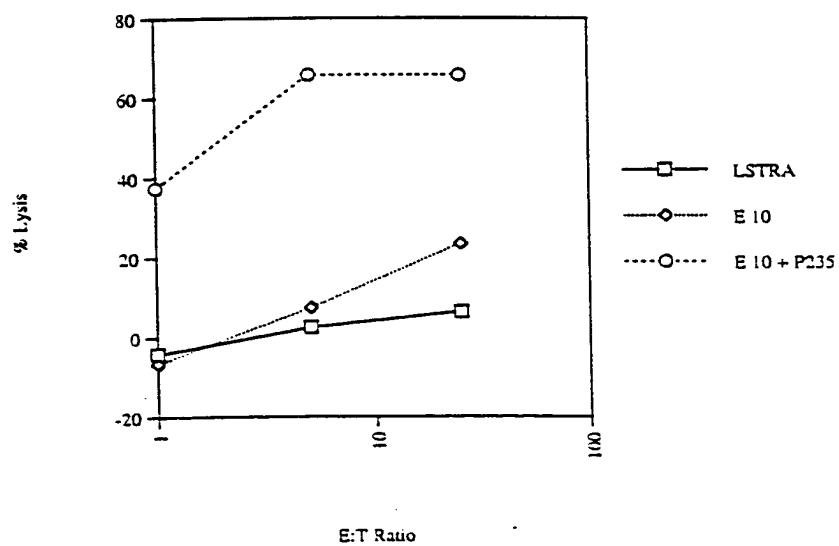


FIG. 6A and 6B

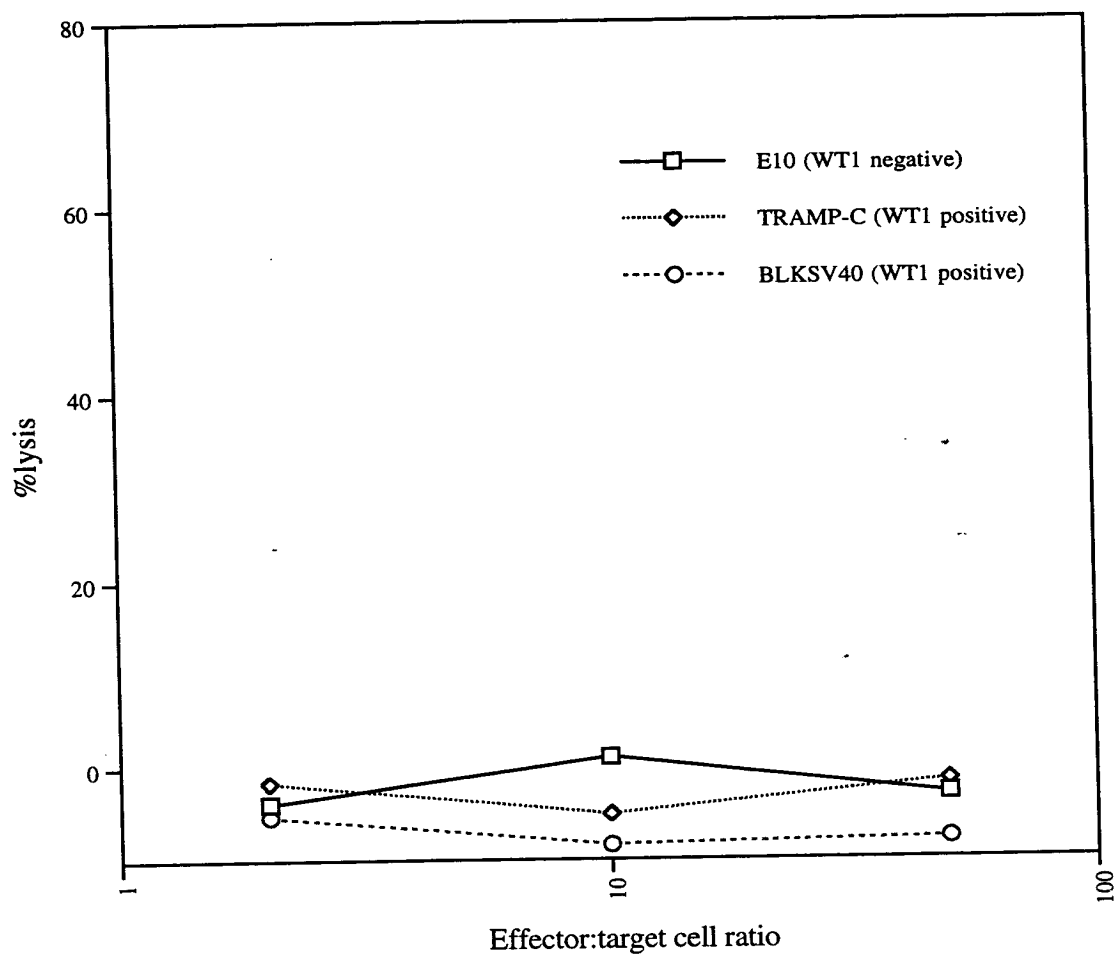


FIG. 7A

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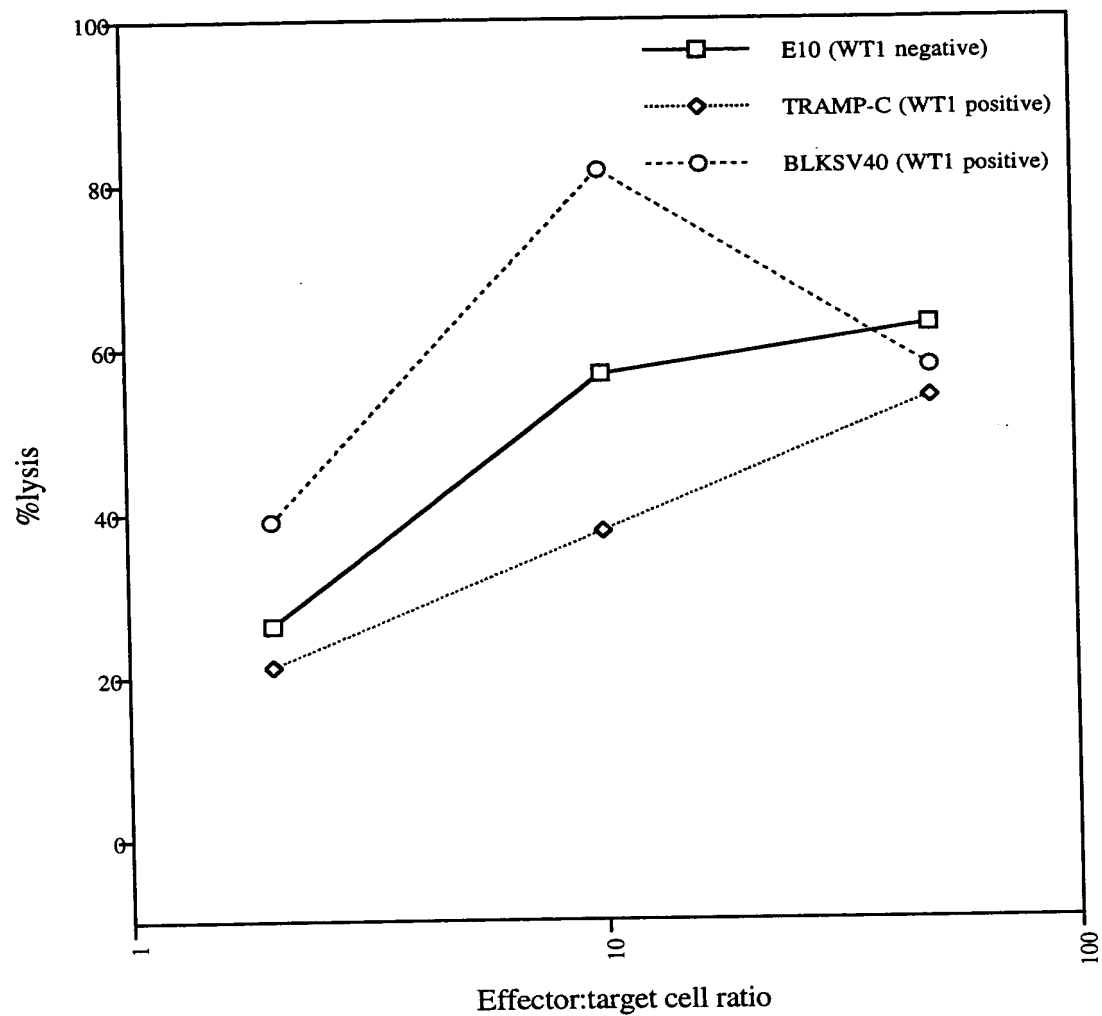


FIG. 7B

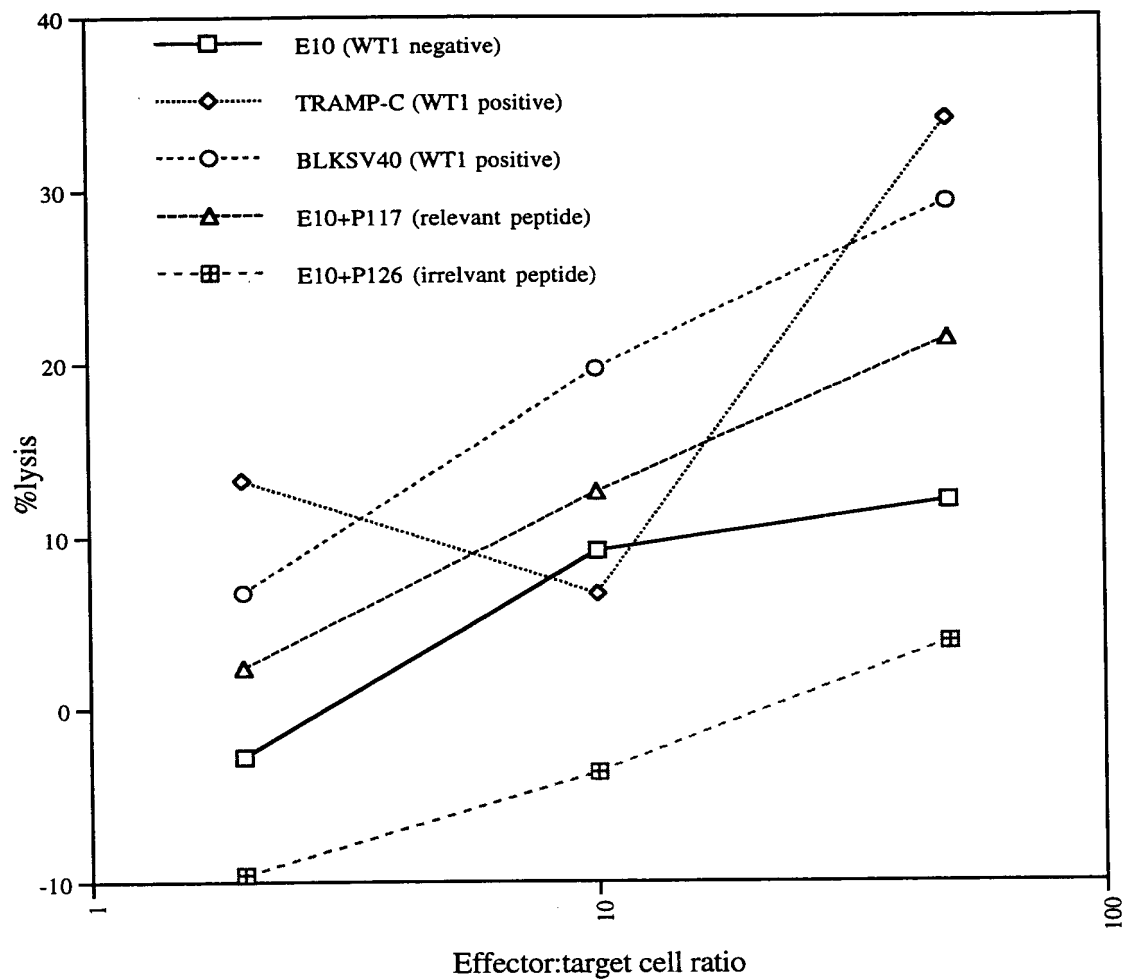


FIG. 7C

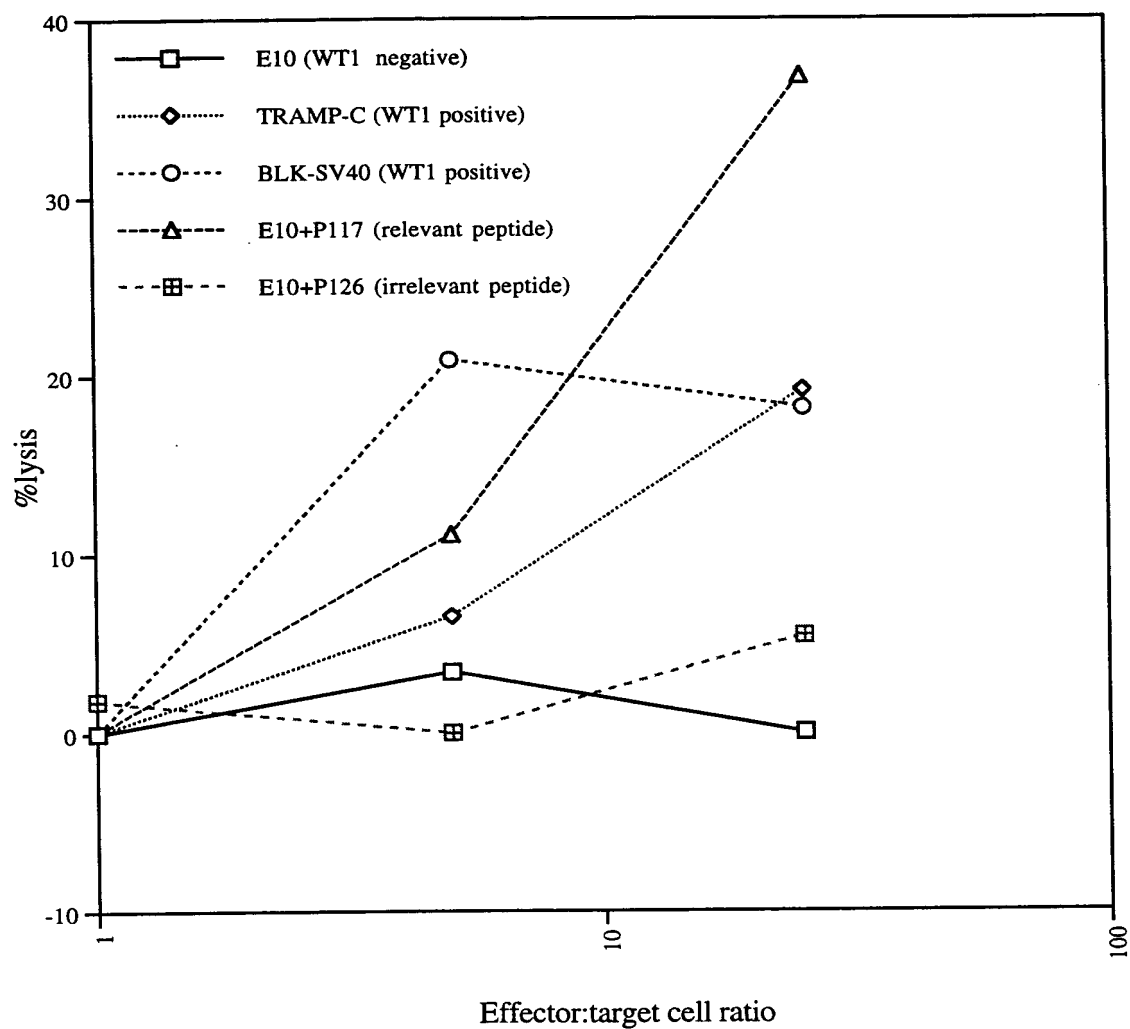


FIG. 7D

09164223 1093098
860660 2249760

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5   10  15  20  25  30  35  40  45  50  55  60  65  70  75
MGSDVRDLNALLPAVPSLGGGGGCGALPVSGAAQWAPVLDFAAPPASAYGSLGGPAPPPAPPPPPPPHPSFIKQE
.....AAAAAAAAAAAAAAAA.....AAAAA.....AAAAAAAAAAAA.....
.....RRRR.....
.....

80  85  90  95  100 105 110 115 120 125 130 135 140 145 150
PSWGGAEPHEEQCLSAFTVHFSGQFTGTAGACRYGPFPPPPSQASSGQARMFPNAPYLPSCLESQPAIRNQYS
.....AAA.....AAAA.....AAA.....AAAAA.....
.....RRRR.....RRRR.....
.....DDDDDDDD.....
.....

155 160 165 170 175 180 185 190 195 200 205 210 215 220 225
TVTFDGTPSYGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPVYGCHTPTDSTGSQALLRTPYSSDN
.....AAAAA.....AAAAA.....AA
.....RRRR.....DDDDDDDDDDDDDD...
.....

230 235 240 245 250 255 260 265 270 275 280 285 290 295 300
LYQMTSQLECMWTWNQMNLGATLKGVAAGSSSVKWTGQSNHSTGYESDNHTTPILCGAQYRIHTHGVRGIQDV
AAAAA.....AAA.AAA.....AAAAAAAAA
.....RRRRRRRRRR.....RRRR.....RRRR.....
DDDDDD.....DDDDDDDDDD.....dddd.....

305 310 315 320 325 330 335 340 345 350 355 360 365 370 375
RRVPGVAPTLVRSASETSEKRPFMCAYPGCNKRYFKLSHLQMHSRKHTGEKPYQCDFKDCERRFSRSDQLKRHQR
AAAAA..AAAAAAAAA.....AAAA.AAAAAAAAAA.
.....RRRR.....RRRR.....
.....DDDDDD.....

380 385 390 395 400 405 410 415 420 425 430 435 440 445 450
RHTGVKPFQCKTCQKRFSDHLKTHTRHTGTSEKPFSCRWPSQCKKFARSDELVRHHNMHQRNMTKLQALAL
.....AAAA.AAAA..AA.....AAAA.....AAA.....AAAAA.....AAA.....
.....RRRR..RRRR.....
.....dddddddddd.....

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FIG. 8A

0016423 093099

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5   10  15  20  25  30  35  40  45  50  55  60  65  70  75
MGSDVRDLNALLPAVSSLGGGGGCGLPVSGAAQWAPVLDFAFPGASAYGSLGGPAPPPAPPPPPPPPHSFIKQE
.....AAAAAAAAAAAAAAAA.....AAAAA.....AAAAAAAAAAAA.....
.....RRRR.....
.....

80  85  90  95  100 105 110 115 120 125 130 135 140 145 150
PSWGGAEPHEEQCLSAFTLHFGQFTGTAGACRYGPFPPPPPSQASSGQARMFPNAPYLPSCLESQPTIRNQYS
.....AAAA.....AAA.....AAAAA.....
.....RRRR.....RRRR.....
.....DDDDDDDD.....

155 160 165 170 175 180 185 190 195 200 205 210 215 220 225
TVTFDGAESYGHTPSHHAAQFPNHSFKHEDPMGQQGSLGEQQYSVPPPYGCHTPTDCTGSQALLLRTPYSSDN
.....AAAA.....AAAAA.....AA
.....RRRR.....DDDDDDDDDDDDDD.....

230 235 240 245 250 255 260 265 270 275 280 285 290 295 300
LYQMTSQLECMTNQMNLGATLKGMAAGSSSVKWTGQSNHGIGYESDNHTAPILCGAQYRIHTGVFRGIQDV
AAAAA.....AAA.AAA.....AAAAAAAAA
.....RRRRRRRRRR.....RRRR.....RRRR.....
DDDDDD.....DDDDDDDDDDDD.....dddd.....

305 310 315 320 325 330 335 340 345 350 355 360 365 370 375
RRVSGVAPTLVRSASETSEKRPFMCAYPGCNKRYFKLSHLQMHRSKHTGEKPYQCDFKDCERRFRSDQLKRHQR
AAAAA.....AAAAA.....AAAA.AAAAAA.....
.....RRRR.....RRRR.....
DDDDDDDDDDDD.....

380 385 390 395 400 405 410 415 420 425 430 435 440 445 450
RHTGVKPFQCKTCQRKFSRSDHLKTHTRTHTGKTSEKPFSCRWHSCQKKFARSDELVRHHNMHQRNMTKLHVAL
.....AAAA.AAAA..AA.....AAAA.....AA.....AAAAA.....AAAA.....
.....RRRR..RRRR.....
.....dddddddddd.....

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FIG. 8B